SEQUENCE LISTING

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<110> Berzofsky, Jay A.
      Pastan, Ira H.
      Terabe, Masaki
      The Government of the United States of America
         as represented by The Secretary of the
         Department of Health and Human Services
<120> Immunogenic Peptides of XAGE-1
<130> 015280-485100US
<140> US 10/582,703
<141> 2006-06-12
<150> US 60/529,025
<151> 2003-12-12
<150> WO PCT/US04/41639
<151> 2004-12-13
<160> 45
<170> PatentIn Ver. 2.1
<210> 1
<211> 246
<212> DNA
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<223> xage-1 p9, 9kD protein expressed from XAGE-1 gene
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atg gag agc ccc aaa aag aag aac cag cag ctg aaa gtc ggg atc cta
                                                                   48
Met Glu Ser Pro Lys Lys Lys Asn Gln Gln Leu Lys Val Gly Ile Leu
                                     10
cac ctg ggc agc aga cag aag atc agg ata cag ctg aga tcc cag
His Leu Gly Ser Arg Gln Lys Lys Ile Arg Ile Gln Leu Arg Ser Gln
             20
tgc gcg aca tgg aag gtg atc tgc aag agc tgc atc agt caa aca ccg
                                                                   144
Cys Ala Thr Trp Lys Val Ile Cys Lys Ser Cys Ile Ser Gln Thr Pro
ggg ata aat ctg gat ttg ggt tcc ggc gtc aag gtg aag ata ata cct
                                                                   192
Gly Ile Asn Leu Asp Leu Gly Ser Gly Val Lys Val Lys Ile Ile Pro
     50
                         55
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aaa gag gaa cac tgt aaa atg cca gaa gca ggt gaa gag caa cca caa'
Lys Glu Glu His Cys Lys Met Pro Glu Ala Gly Glu Glu Gln Pro Gln
gtt taa
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Val
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Met Glu Ser Pro Lys Lys Lys Asn Gln Gln Leu Lys Val Gly Ile Leu
His Leu Gly Ser Arg Gln Lys Lys Ile Arg Ile Gln Leu Arg Ser Gln
             20
Cys Ala Thr Trp Lys Val Ile Cys Lys Ser Cys Ile Ser Gln Thr Pro
Gly Ile Asn Leu Asp Leu Gly Ser Gly Val Lys Val Lys Ile Ile Pro
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Lys Glu Glu His Cys Lys Met Pro Glu Ala Gly Glu Glu Gln Pro Gln
 65
                     70
Val
<210> 3
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<223> xage-1 pl6, 16.3 kD protein expressed from XAGE-1
      gene
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<222> (1)..(441)
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atg ctc ctt tgg tgc cca cct cag tgc gca tgt tca ctg ggc gtc ttc
                                                                   48
Met Leu Leu Trp Cys Pro Pro Gln Cys Ala Cys Ser Leu Gly Val Phe
 1
cca tcg gcc cct tcg cca gtg tgg gga acg cgg cgg agc tgt gag ccg
                                                                   96
Pro Ser Ala Pro Ser Pro Val Trp Gly Thr Arg Arg Ser Cys Glu Pro
             20
                                 25
gcg act cgg gtc cct gag gtc tgg att ctt tct ccg cta ctg aga cac
Ala Thr Arg Val Pro Glu Val Trp Ile Leu Ser Pro Leu Leu Arg His
                             40
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Gry			aca Thr													192
gta Val 65																240
cta Leu																288
cag Gln					-	_		_	_	-	_		_			336
ccg Pro																384
cct Pro																432
caa Gln 145	_	taa														441
<210	> 4 > 14	16														
<212 <213			sapie	ens												
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Pro Lys Glu Glu His Cys Lys Met Pro Glu Ala Gly Glu Glu Gln Pro
    130
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Gln Val
145
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      peptide derived from xage-1 14
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<222> (1)
<223> Xaa = any amino acid (X-1)
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<222> (2)
\langle 223 \rangle Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
<220>
<221> MOD RES
<222> (3)
<223> Xaa = a \ hydrophobic \ residue, \ Met \ or \ Ala \ (X-3)
<220>
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<222> (10)
\langle 223 \rangle Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
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Xaa Xaa Xaa Pro Ser Ala Pro Ser Pro Xaa
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<210> 6
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<213> Artificial Sequence
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<223> Description of Artificial Sequence:xage-1 14,
      immunogenic amino terminal end of xage-1, xage-1
      residues 14-23
Gly Val Phe Pro Ser Ala Pro Ser Pro Val
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: 1Y xage-1 14,
      variant of xage-1 14, immunogenic peptide derived
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Tyr Val Phe Pro Ser Ala Pro Ser Pro Val
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<210> 8
<211> 10
<212> PRT
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<223> Description of Artificial Sequence: 2L xage-1 14,
      variant of xage-1 14, immunogenic peptide derived
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<400> 8
Gly Leu Phe Pro Ser Ala Pro Ser Pro Val
                  5
<210> 9
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: 3M xage-1 14,
      variant of xage-1 14, immunogenic peptide derived
      from xage-1 14
<400> 9
Gly Val Met Pro Ser Ala Pro Ser Pro Val
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<210> 10
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: 1Y2L xage-1 14,
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Tyr Leu Phe Pro Ser Ala Pro Ser Pro Val
                  5
<210> 11
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Gly Leu Met Pro Ser Ala Pro Ser Pro Val
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<223> Description of Artificial Sequence:modified xage-1
      14 peptide, immunogenic peptide derived from
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Gly Val Trp Pro Ser Ala Pro Ser Pro Val
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      xage-1 14
<400> 13
Gly Val Tyr Pro Ser Ala Pro Ser Pro Val
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<210> 14
<211> 10
<212> PRT
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Thr Val Trp Pro Ser Ala Pro Ser Pro Met
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<210> 15
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<213> Artificial Sequence
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      14 peptide, immunogenic peptide derived from
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Ser Met Tyr Pro Ser Ala Pro Ser Pro Ile
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      14 peptide, immunogenic peptide derived from
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Ser Val Phe Pro Ser Ala Pro Ser Pro Thr
<210> 17
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Gly Val Trp Pro Ser Ala Pro Ser Pro Met
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<210> 18
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<210> 19
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Gly Leu Trp Pro Ser Ala Pro Ser Pro Val
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Ile Val Trp Pro Ser Ala Pro Ser Pro Val
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<400> 21
Gly Leu Ala Pro Ser Ala Pro Ser Pro Val
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<400> 22
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<210> 23
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<220>
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      14 peptide, immunogenic peptide derived from
      xage-1 14
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Tyr Leu Phe Pro Ser Ala Pro Ser Pro Met
                  5
<210> 24
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      xage-1 14
<400> 24
Tyr Leu Ala Pro Ser Ala Pro Ser Pro Ile
                  5
<210> 25
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<213> Artificial Sequence
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<400> 25
Tyr Leu Ala Pro Ser Ala Pro Ser Pro Val
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                 5
<210> 26
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<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence:nucleic acid
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<400> 26
ggcgtcttcc catcggcccc ttcgccagtg
                                                                   30
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<211> 30
<212> DNA
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<223> Description of Artificial Sequence:nucleic acid
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<400> ggcgto	27 catgo catoggocoo ttogocagtg	30
<210> <211> <212> <213>	30	
<220> <223>	Description of Artificial Sequence: nucleic acid sequence encoding SEQ ID NO:11 preferred form	
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<210> <211> <212> <213>	30	
<220> <223>	Description of Artificial Sequence: nucleic acid sequence encoding SEQ ID NO:11 preferred form	
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<210><211><212><212><213>	30	
<220> <223>	Description of Artificial Sequence:nucleic acid sequence encoding SEQ ID NO:11 preferred form	
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<220> <223>	Description of Artificial Sequence:nucleic acid sequence encoding SEQ ID NO:11 preferred form	
<400> ggcctc	31 gatgc catcggcccc ttcgccagtg	30
<210><211><211><212><213>	10	

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<223> Description of Artificial Sequence:xage-1 33,
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Ala Thr Arg Val Pro Glu Val Trp Ile Leu
<210> 33
<211> 10
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:xage-1 57,
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His Thr Ala Ser Pro Arg Ser Pro Val Met
<210> 34
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<222> (2)
\langle 223 \rangle Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
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\langle 223 \rangle Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD RES
<222> (10)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
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                  5
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<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:immunogenic
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<221> MOD RES
<222> (1)
<223> Xaa = any amino acid (X-1)
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<221> MOD_RES
<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
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<221> MOD RES
<222> (10)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
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Xaa Leu Xaa Pro Ser Ala Pro Ser Pro Xaa
                   5
<210> 36
<211> 10
<212> PRT
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<223> Description of Artificial Sequence:immunogenic
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<221> MOD RES
<222> (2)
\langle 223 \rangle Xaa = Leu, Met, Ala, Ile, Val or Thr \langle X-2 \rangle
<220>
<221> MOD RES
<222> (10)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
<400> 36
Xaa Xaa Met Pro Ser Ala Pro Ser Pro Xaa
                   5
<210> 37
<211> 10
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence:immunogenic
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<221> MOD RES
<222> (1)
<223> Xaa = any amino acid (X-1)
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<221> MOD_RES
<222> (2)
<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
<220>
<221> MOD RES
<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
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<223> Xaa = any amino acid (X-1)
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\langle 223 \rangle Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
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<221> MOD RES
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<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD_RES
<222> (9)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
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Xaa Xaa Xaa Pro Ser Ala Pro Ser Xaa
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<210> 39
<211> 9
<212> PRT
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<223> Xaa = any amino acid (X-1)
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<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
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<221> MOD RES
<222> (3)
<223> Xaa = a \ hydrophobic \ residue, \ Met \ or \ Ala \ (X-3)
<220>
<221> MOD RES
<222> (9)
\langle 223 \rangle Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
<400> 39
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                   5
<210> 40
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<212> PRT
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<223> Description of Artificial Sequence: 9-mer created
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<221> MOD RES
<222> (2)
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<220>
<221> MOD RES
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<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD RES
<222> (9)
\langle 223 \rangle Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
Xaa Xaa Xaa Pro Ser Ala Ser Pro Xaa
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<210> 41
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence:overall formula
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<222> (1)
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<221> MOD RES
<222> (2)
<223> Xaa = Leu, Met, Ala, Ile, Val or Thr (X-2)
<220>
<221> MOD RES
<222> (3)
<223> Xaa = a hydrophobic residue, Met or Ala (X-3)
<220>
<221> MOD RES
<222> (7)
<223> Xaa = Pro or absent (X-5), when absent, X-6 is Ser
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<221> MOD RES
<222> (8)
<223> Xaa = Ser or absent (X-6), when absent, X-5 and X-7 are Pro
<220>
<221> MOD RES
<222> (9)
<223> Xaa = Pro or absent (X-7), when absent, X-5 is Pro and X-6
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<222> (10)
<223> Xaa = Val, Met, Leu, Ala, Ile or Thr (X-4)
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Xaa Xaa Xaa Pro Ser Ala Xaa Xaa Xaa
                  5
<210> 42
<211> 27
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<223> Description of Artificial Sequence:exemplar
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                                                                   27
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<210> 43
<211> 27
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence:exemplar
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ggcgtcttcc catcggcccc tccagtg
                                                                   27
<210> 44
<211> 27
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: exemplar
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<400> 44
ggcgtcttcc catcggcctc gccagtg
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<210> 45
<211> 637
<212> DNA
<213> Homo sapiens
<220>
<223> complete nucleic acid sequence of XAGE-1 with
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ccacagecet taaggeacga gggaacetea etgegeatge teetttggtg eccaceteag 120
tgcgcatgtt cactgggcgt cttcccatcg gccccttcgc cagtgtgggg aacgcggcgg 180
agetgtgage eggegaeteg ggteeetgag gtetggatte ttteteeget aetgagaeae 240
ggcggacaca cacaaacaca gaaccacaca gccagtccca ggagcccagt aatggagagc 300
cccaaaaaga agaaccagca gctgaaagtc gggatcctac acctgggcag cagacagaag 360
aagatcagga tacagctgag atcccagtgc gcgacatgga aggtgatctg caagagctgc 420
atcagtcaaa caccggggat aaatctggat ttgggttccg gcgtcaaggt gaagataata 480
cctaaagagg aacactgtaa aatgccagaa qcaqqtqaaq aqcaaccaca agtttaaatg 540
aagacaagct gaaacaacgc aagctggttt tatattagat atttgactta aactatctca 600
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